

006121" 05514250

DIFFERENTIAL GENE EXPRESSION IN NORMAL TOTAL BRAIN VS. CORPUS CALLOSUM

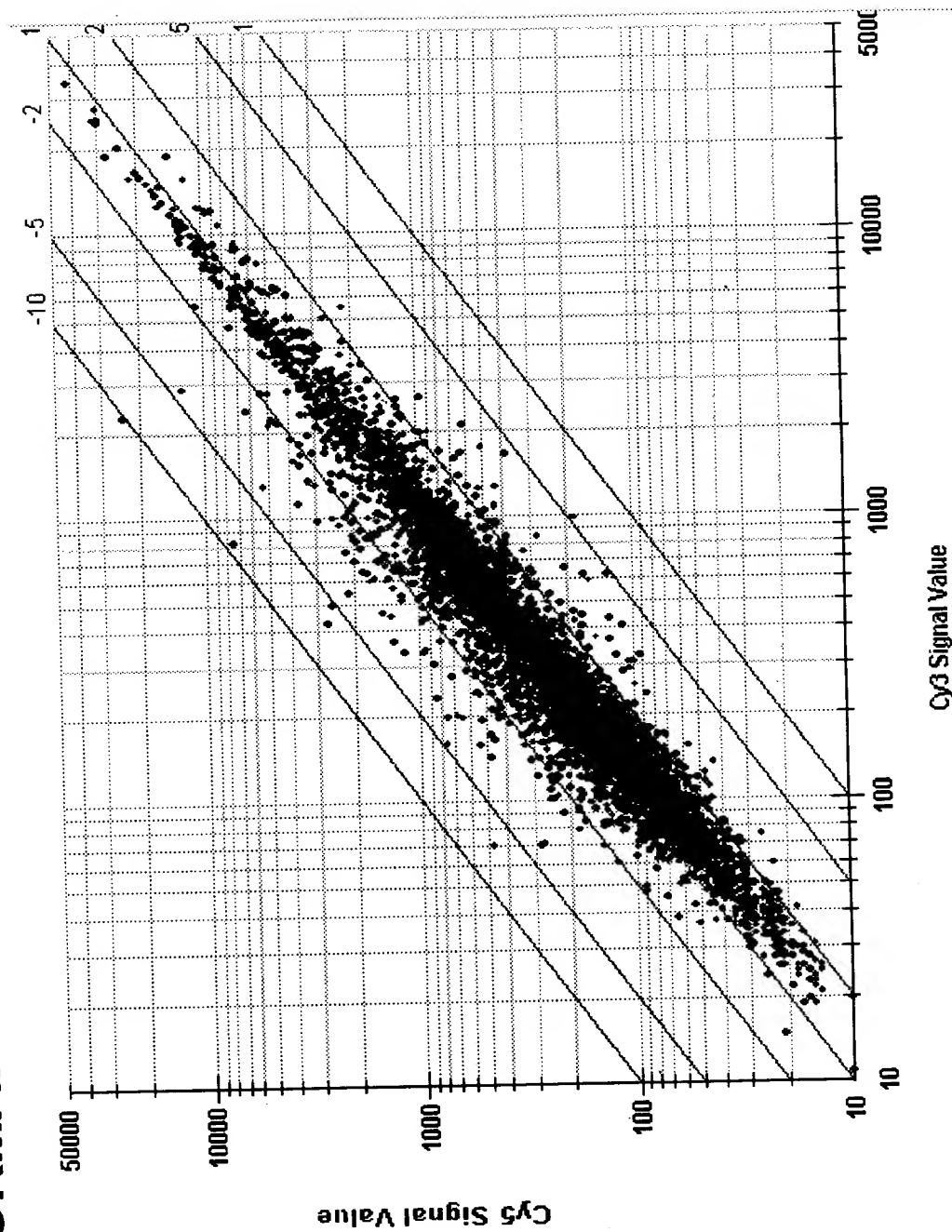


Figure 1A

006T2T" 055F4260

DIFFERENTIAL GENE EXPRESSION IN GLIOBLASTOMA #22 VS. NORMAL BRAIN

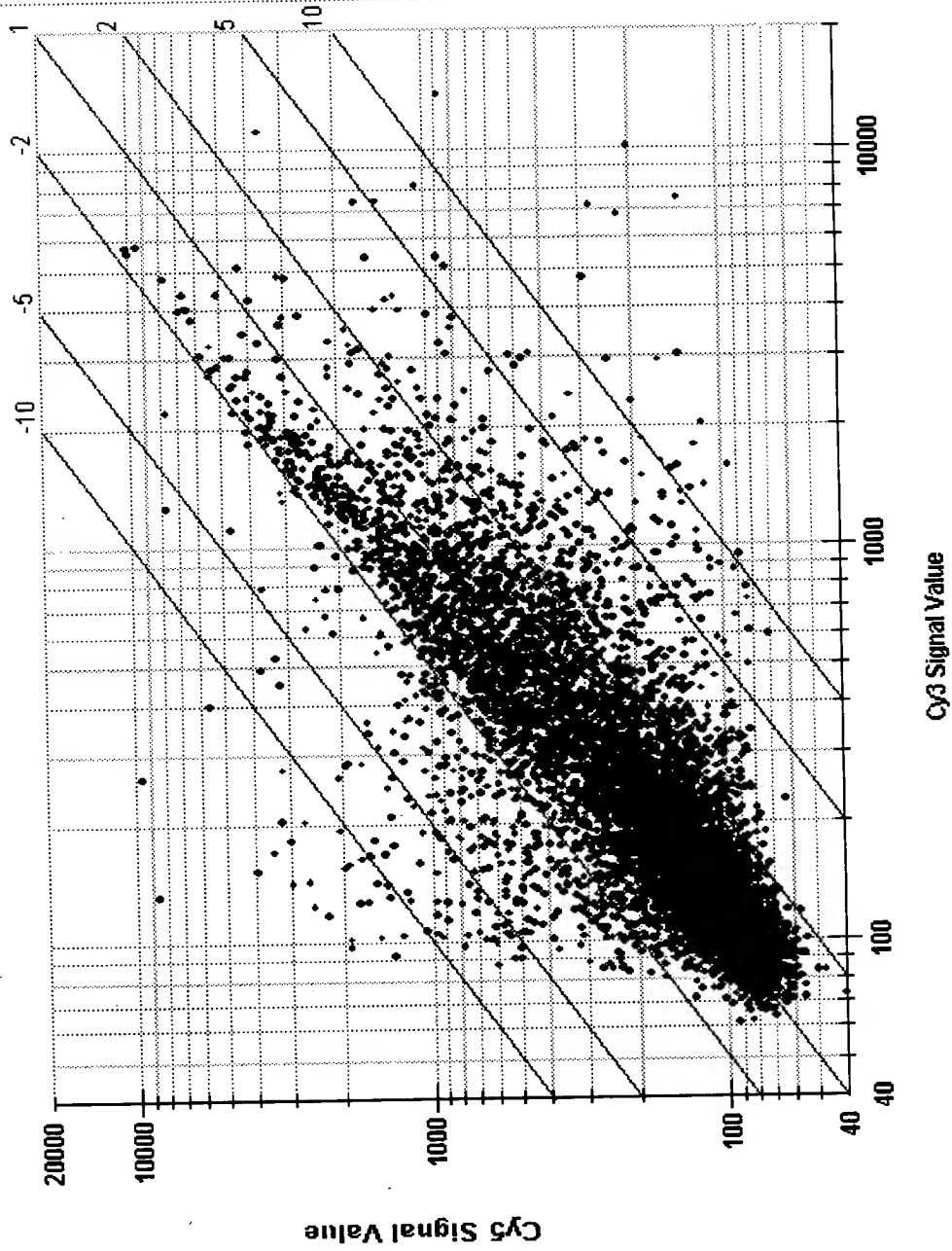


Figure 1B

006T2T" 055T460

39t

Cy3/Cy5 Graph

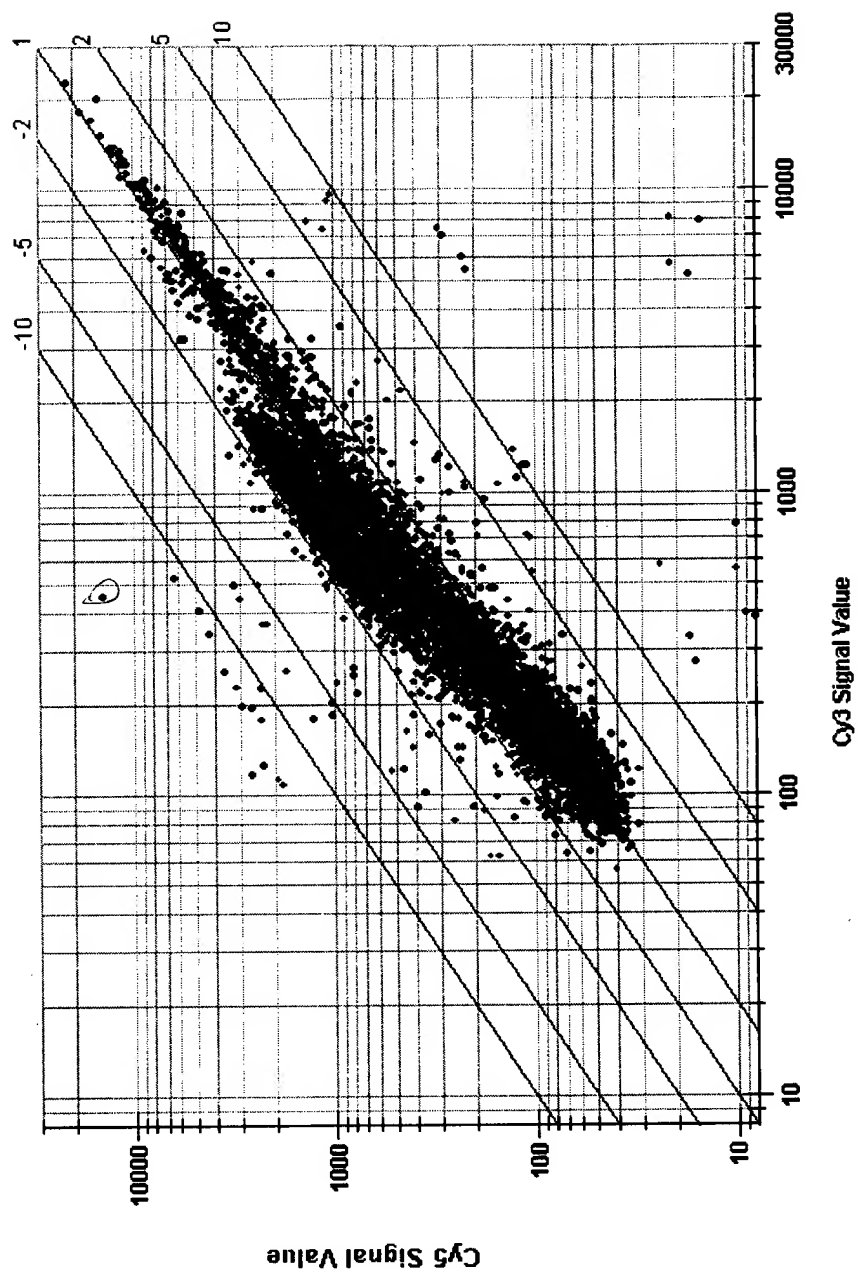


Figure 1C

006T2T"055T4260

39a

Cy3/Cy5 Graph

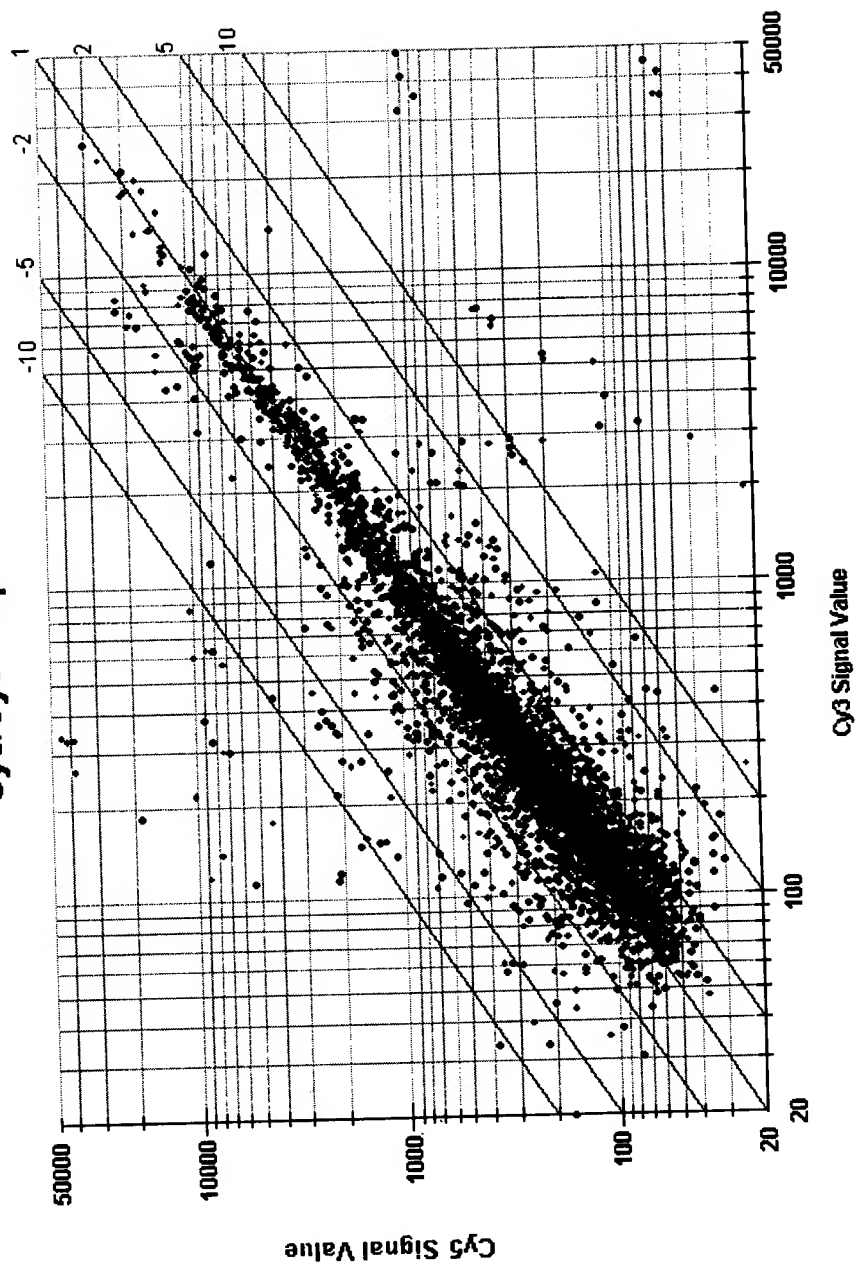


Figure 1D

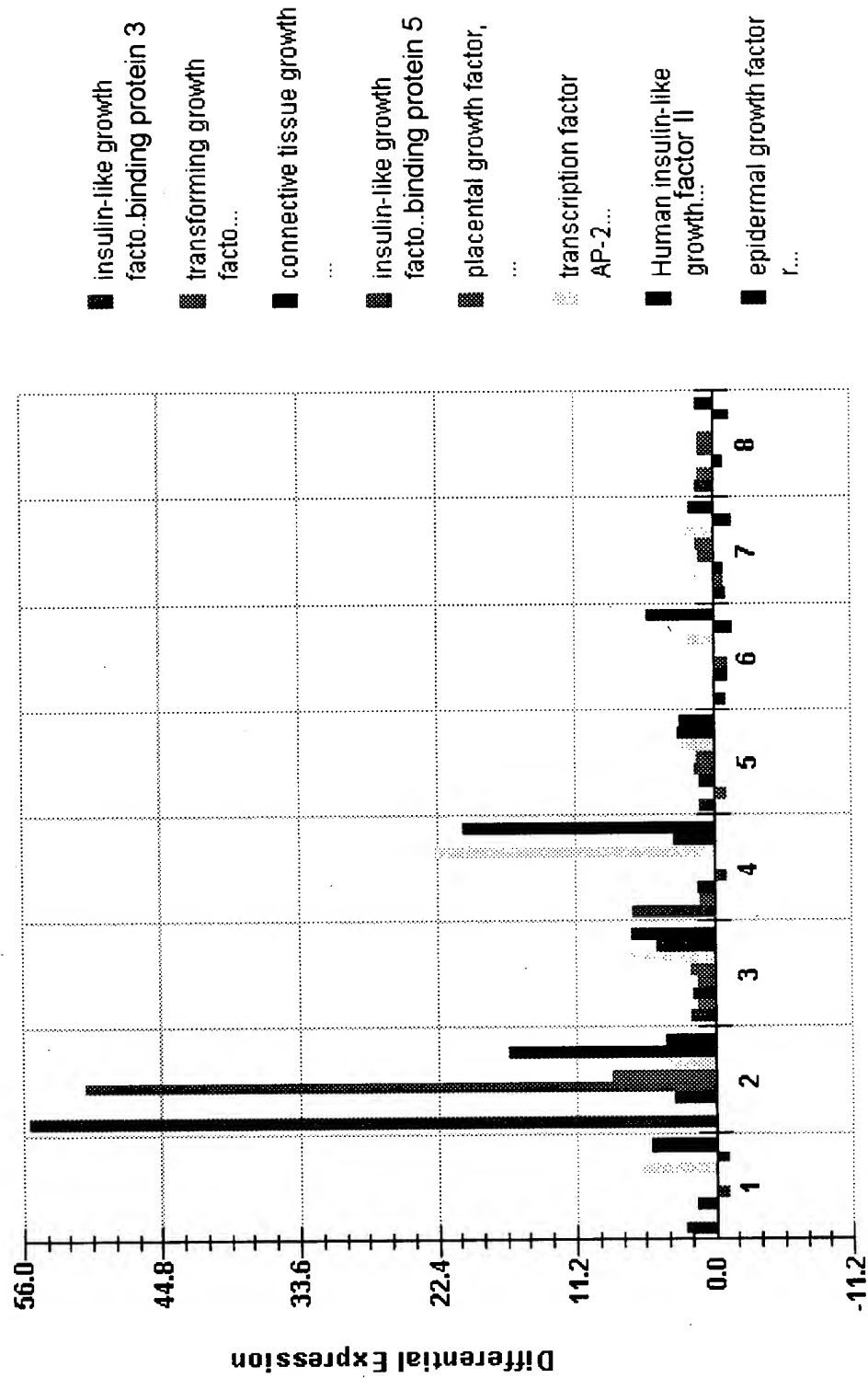


Figure 2

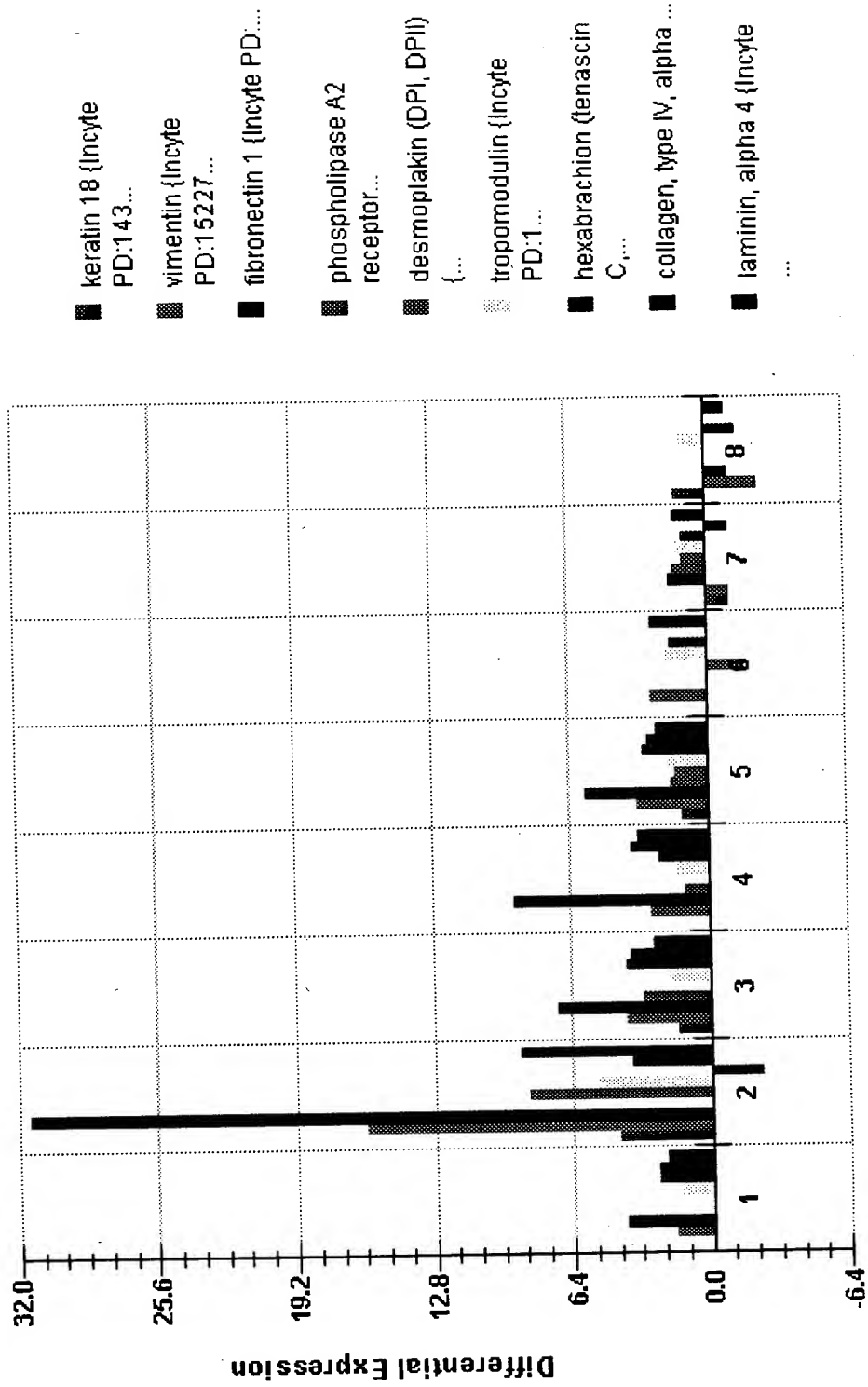


Figure 3

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Growth factors

Series Plot

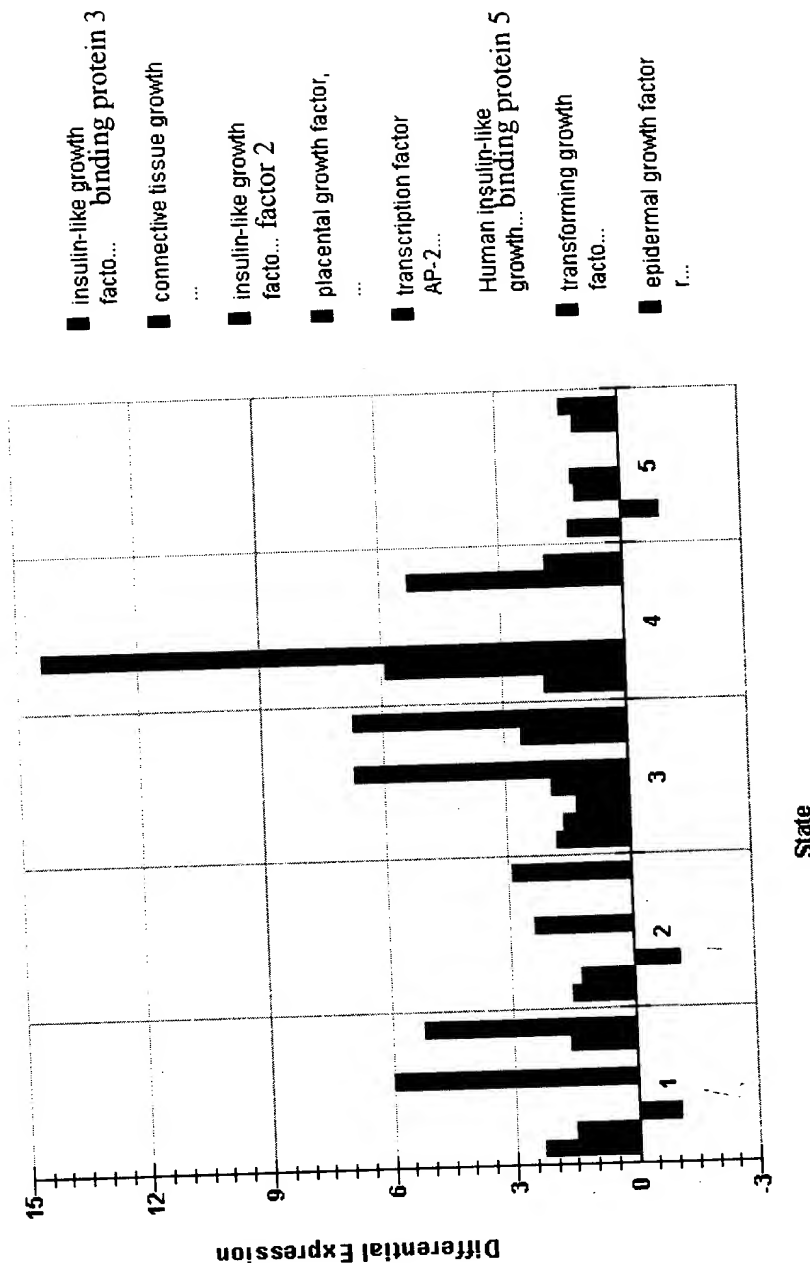


Figure 4A

Matrix group

Series Plot

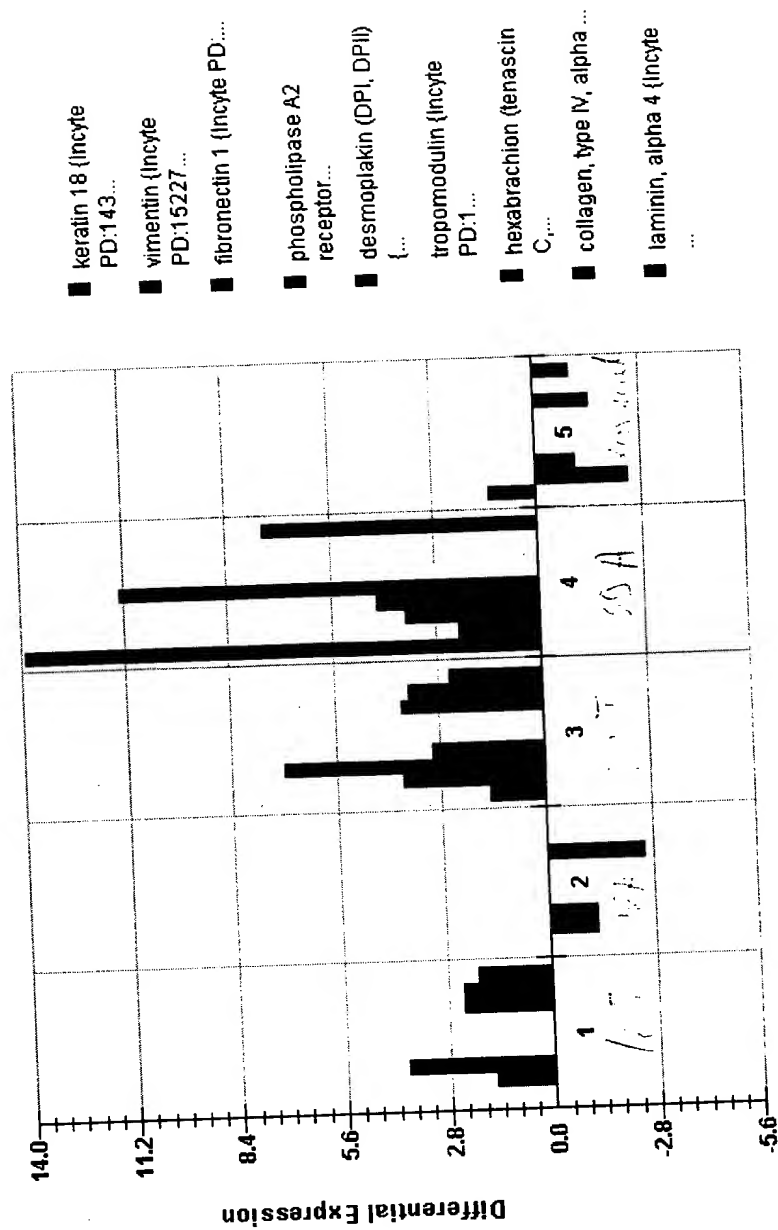


Figure 4B

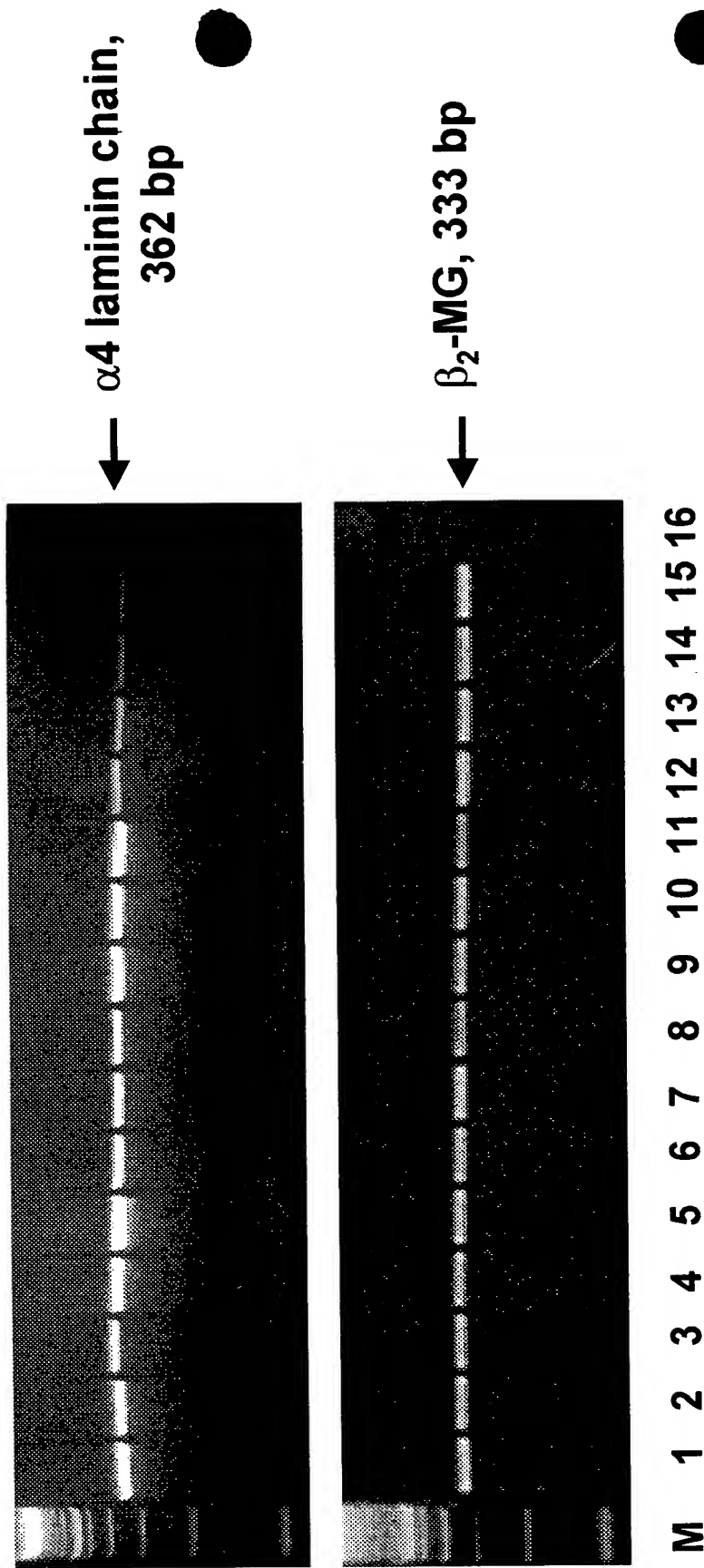


Figure 5 Semiquantitative RT-PCR analysis of expression in brain tumors. Top, expression of the 362 bp gene of $\alpha 4$ laminin chain; bottom, expression of the 333 bp fragment of β_2 -microglobulin gene. Lanes are as follows: 1, GBM* #16; primary tumor; 2, adjacent tissue to the GBM#16*; 3, GBM #22; 4, GBM #39*, primary tumor; 5, adjacent tissue to the GBM #39*; 6, GBM #45; 7, GBM #50; 8, GBM #47; 9, GBM #25, primary tumor; 10, adjacent tissue to the GBM #25; 11, astrocytoma grade II #34; 12, meningioma (benign tumor) #38; 13, normal brain #46; 14, normal brain #40; 15, corpus callosum; 16, control without RT; M, 100 bp DNA ladder.

Figure 5

005T2T" 055T460

γ 1 chain

β 2 chain

β 1 chain

α 4 chain

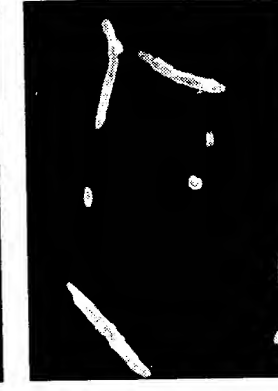
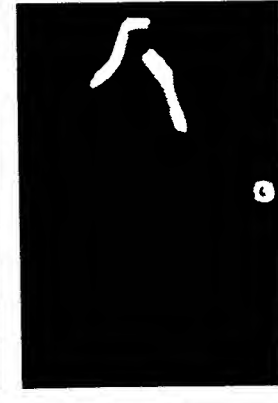
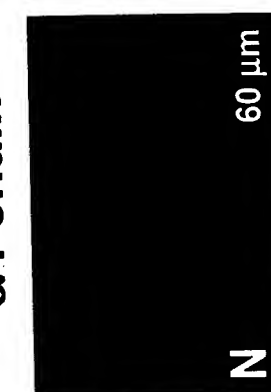
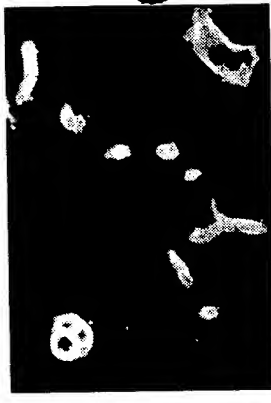


Figure 6

